OIPE

ENTERED

Input Set : A:\SEQLISTCORRECTED2.TXT
Output Set: N:\CRF4\11132002\J073644C.raw

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 3 <110> APPLICANT: Hudson, Debra
 4        van de Winkel, Jan
 5        van Dijk, Marc
 7 <120> TITLE OF INVENTION: HUMAN MONOCLONAL ANTIBODIES TO FC ALPHA
 8        RECEPTOR (CD89)
10 <130> FILE REFERENCE: MXI-211
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C--> 12 **<140> CURRENT APPLICATION NUMBER: US/10/073,644C**
C--> 12 **<141> CURRENT FILING DATE: 2002-02-11**
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12 <150> PRIOR APPLICATION NUMBER: US 60/338,956
13 <151> PRIOR FILING DATE: 2001-11-05
15 <150> PRIOR APPLICATION NUMBER: US 60/268,075
16 <151> PRIOR FILING DATE: 2001-02-12
18 <160> NUMBER OF SEQ ID NOS: 8
20 <170> SOFTWARE: FastSEQ for Windows Version 4.0
22 <210> SEQ ID NO: 1
23 <211> LENGTH: 357
24 <212> TYPE: DNA
25 <213> ORGANISM: Homo sapiens
27 <400> SEQUENCE: 1
28 caggtgcaac tggtggagtc tggggggaggc gtggtccagc ctgggaggtc cctgagactc 60
29 tcctgtgcag cctctggatt caccttcagt agttatgttc tgcactgggt ccgccaggct 120
30 ccaggcaagg ggctggattg ggtggcagtg atatcagatg atggaaggaa taaatacttc 180
31 gcagactccg tgaagggccg attcaccatc tccagagaca attccaagaa cacgctgtat 240
32 ctgcaaatga acagcctgag agctgaggac acggctgtgt attactgtgt gagagaaggg 300
33 tatagcggca gctggtttga ctactggggc cagggaaccc tggtcaccgt ctcctca   357
35 <210> SEQ ID NO: 2
36 <211> LENGTH: 119
37 <212> TYPE: PRT
38 <213> ORGANISM: Homo sapiens
40 <400> SEQUENCE: 2
41 Gln Val Gln Leu Val Glu Ser Gly Gly Gly Val Val Gln Pro Gly Arg
42 1               5                   10                  15
43 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Ser Tyr
44             20                  25                  30
45 Val Leu His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Asp Trp Val
46         35                  40                  45
47 Ala Val Ile Ser Asp Asp Gly Arg Asn Lys Tyr Phe Ala Asp Ser Val
48     50                  55                  60
49 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
50 65                  70                  75                  80
51 Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
52             85                  90                  95
53 Val Arg Glu Gly Tyr Ser Gly Ser Trp Phe Asp Tyr Trp Gly Gln Gly
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RAW SEQUENCE LISTING   DATE: 11/13/2002
PATENT APPLICATION: **US/10/073,644C**   TIME: 14:09:54

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54            100           105           110
55 Thr Leu Val Thr Val Ser Ser
56         115
59 <210> SEQ ID NO: 3
60 <211> LENGTH: 321
61 <212> TYPE: DNA
62 <213> ORGANISM: Homo sapiens
64 <400> SEQUENCE: 3
65 gccatccagt tgacccagtc tccatcctcc ctgtctgcat ctgtaggaga cagagtcacc 60
66 atcacttgcc gggcaagtca gggcattagc agtgctttag cctggtatca gcagaaacca 120
67 gggaaagctc ctaagctcct gatctatggt gcctccagtt tggaaggtgg ggtcccatca 180
68 aggttcagcg gcagtggatc tgggacagat ttcactctca ccatcagcag cctgcagcct 240
69 gaagattttg caacttatta ctgtcaacag tttaatagtt acccattcac tttcggccct 300
70 gggaccaaag tggatatcaa a                                       321
72 <210> SEQ ID NO: 4
73 <211> LENGTH: 107
74 <212> TYPE: PRT
75 <213> ORGANISM: Homo sapiens
77 <400> SEQUENCE: 4
78 Ala Ile Gln Leu Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly
79 1             5               10              15
80 Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Gly Ile Ser Ser Ala
81           20              25              30
82 Leu Ala Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu Ile
83         35              40              45
84 Tyr Gly Ala Ser Ser Leu Glu Gly Gly Val Pro Ser Arg Phe Ser Gly
85     50              55              60
86 Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln Pro
87 65              70              75              80
88 Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Phe Asn Ser Tyr Pro Phe
89               85              90              95
90 Thr Phe Gly Pro Gly Thr Lys Val Asp Ile Lys
91           100             105
92 <210> SEQ ID NO: 5
93 <211> LENGTH: 357
94 <212> TYPE: DNA
95 <213> ORGANISM: Homo sapiens
97 <400> SEQUENCE: 5
98 caggtgcagc tggtggagtc tgggggaggc gtggtccagc ctgggaggtc cctgagactc 60
99 tcctgtgcag cctctggatt caccttcagt agctatgcta tgcactgggt ccgccaggct 120
100 ccaggcaagg ggctggagtg ggtggcagtt atatcatatg atggaagaaa taaagactac 180
101 gcagactccg tgaagggccg attcaccatc tccagagaca attccaagaa cacgctgtat 240
102 ctgcaaatga acagcctgag agctgaggac acggctgtgc attactgtgc gaggcttgac 300
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105 <210> SEQ ID NO: 6
106 <211> LENGTH: 119
107 <212> TYPE: PRT
108 <213> ORGANISM: Homo sapiens
110 <400> SEQUENCE: 6
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111 Gln Val Gln Leu Val Glu Ser Gly Gly Gly Val Val Gln Pro Gly Arg
112 1               5               10              15
113 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Ser Tyr
114             20              25              30
115 Ala Met His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
116         35              40              45
117 Ala Val Ile Ser Tyr Asp Gly Arg Asn Lys Asp Tyr Ala Asp Ser Val
118     50              55              60
119 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
120 65              70              75              80
121 Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val His Tyr Cys
122             85              90              95
123 Ala Arg Leu Asp Trp Gly Tyr Asp Ala Phe Asp Ile Trp Gly Gln Gly
124             100             105             110
125 Thr Met Val Thr Val Ser Ser
126         115
128 <210> SEQ ID NO: 7
129 <211> LENGTH: 327
130 <212> TYPE: DNA
131 <213> ORGANISM: Homo sapiens
133 <400> SEQUENCE: 7
134 gaaattgtgt tgacgcagtc tccaggcacc ctgtctttgt ctccagggga aagagccacc 60
135 ctctcctgca gggccagtca gagtgttagc agcagctact tagcctggta ccagcagaag 120
136 cctggccagg ctcccaggct cctcatctat ggtgcatcca gcagggccac tggcatccca 180
137 gacaggttca gtggcagtgg gtctgggaca gacttcactc tcaccatcag cagactggag 240
138 cctgaagatt ttgcagtgta ttactgtcag cagtatggta gctcacctcc gtacactttt 300
139 ggccagggga ccaagctgga gatcaaa 327
141 <210> SEQ ID NO: 8
142 <211> LENGTH: 109
143 <212> TYPE: PRT
144 <213> ORGANISM: Homo sapiens
146 <400> SEQUENCE: 8
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148 1               5               10              15
149 Glu Arg Ala Thr Leu Ser Cys Arg Ala Ser Gln Ser Val Ser Ser Ser
150             20              25              30
151 Tyr Leu Ala Trp Tyr Gln Gln Lys Pro Gly Gln Ala Pro Arg Leu Leu
152         35              40              45
153 Ile Tyr Gly Ala Ser Ser Arg Ala Thr Gly Ile Pro Asp Arg Phe Ser
154     50              55              60
155 Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Arg Leu Glu
156 65              70              75              80
157 Pro Glu Asp Phe Ala Val Tyr Tyr Cys Gln Gln Tyr Gly Ser Ser Pro
158             85              90              95
159 Pro Tyr Thr Phe Gly Gln Gly Thr Lys Leu Glu Ile Lys
160         100             105
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VERIFICATION SUMMARY
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L:12 M:270 C: Current Application Number differs, Replaced Current Application No
L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date